

Amendments to the Specification

Please amend the title of the application as follows:

***METHODS AND COMPOSITIONS FOR THE
DETECTION OF CERVICAL CANCER***

Please amend the paragraph bridging pages 46 and 47 of the application and previously amended on July 30, 2001, as follows:

Ten masses were detected by mass spectrometry from seven of the CvC-3H peaks. Amino acid sequence was obtained for three peptides, two by Edman degradation and one by carboxypeptidase-MALDI-TOF analysis. The sequences obtained for these peptides, shown in Table 4, match a protein known as IEF SSP 9502 or "novel human nuclear phosphoprotein" (Honore *et al.* (1994) *supra*; GenBank Accession #L07758). ~~The nucleotide sequence of the cDNA encoding IEF SSP 9502 is shown in SEQ ID NO: 47, and~~ The complete amino acid sequence for this protein, as derived from the gene sequence, is shown in SEQ ID NO: 10 ~~and SEQ ID NO: 48~~. Seven other masses from peak fractions separated on the CvC-3H tryptic map also matched those of predicted tryptic fragments from this protein. Mass correlation data of tryptic peptides from CvC-3H are summarized in Table 4. The predicted molecular weight of the nuclear phosphoprotein, based upon its nucleotide sequence, is 55 kDa, whereas its observed molecular weight by 2-D gel analysis is 79 kDa (Honore *et al.* (1994) *supra*).

Please delete from the application the Sequence Listing filed on July 30, 2001, and replace it with the attached pages marked "Sequence Listing." A computer-readable copy of the amended sequence listing is enclosed.